## SEQUENCE LISTING

- (2) INFORMATION FOR SEQ ID NO:1:
  - (i) SEQUENCE CHARACTERISTICS:
    - (A) LENGTH: 65 amino acids
    - (B) TYPE: amino acid(D) TOPOLOGY: linear
  - (ii) MOLECULE TYPE: peptide
  - (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 1:

Val Val Tyr Thr Asp Cys Thr Glu Ser Gly Gln Asn Leu Cys Leu Cys 1 5 10 15

Glu Gly Ser Asn Val Cys Gly Gln Gly Asn Lys Cys Ile Leu Gly Ser 20 25 30

Asp Gly Glu Lys Asn Gln Cys Val Thr Gly Glu Gly Thr Pro Lys Pro
35 40 45

Gln Ser His Asn Asp Gly Asp Phe Glu Glu Ile Pro Glu Glu Tyr Leu 50 60

Gln 65

- (2) INFORMATION FOR SEQ ID NO:2:
  - (i) SEQUENCE CHARACTERISTICS:
    - (A) LENGTH: 65 amino acids
    - (B) TYPE: amino acid
    - (D) TOPOLOGY: linear
  - (ii) MOLECULE TYPE: peptide
  - (xi) SEQUENCE DESCRIPTION: SEO ID NO:2:

Ile Thr Tyr Thr Asp Cys Thr Glu Ser Gly Gln Asn Leu Cys Leu Cys
1 5 15

Glu Gly Ser Asn Val Cys Gly Lys Gly Asn Lys Cys Ile Leu Gly Ser 20 25 30

Asn Gly Lys Gly Asn Gln Cys Val Thr Gly Glu Gly Thr Pro Asn Pro
35 40 45

Glu Ser His Asn Asn Gly Asp Phe Glu Glu Ile Pro Glu Glu Tyr Leu
50 60

Gln 65

## (2) INFORMATION FOR SEQ ID NO:3:

- (i) SEQUENCE CHARACTERISTICS:
  - (A) LENGTH: 65 amino acids
  - (B) TYPE: amino acid
  - (D) TOPOLOGY: linear
- (ii) MOLECULE TYPE: peptide
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:3:

Val Val Tyr Thr Asp Cys Thr Glu Ser Gly Gln Asn Leu Cys Leu Cy 1 5 10 15

Glu Gly Ser Asn Val Cys Gly Gln Gly Asn Lys Cys Ile Leu Gly Ser 20 25 30

Gln Gly Lys Asp Asn Gln Cys Val Thr Gly Glu Gly Thr Pro Lys Pro
35 40 45

Gln Ser His Asn Gln Gly Asp Phe Glu Pro Ile Pro Glu Asp Ala Asp 50 55

Glu 65

## (2) INFORMATION FOR SEQ ID NO:4:

- (i) SEQUENCE CHARACTERISTICS:
  - (A) LENGTH: 64 amino acids
  - (B) TYPE: amino acid
  - (C) STRANDEDNESS: single
  - (D) TOPOLOGY: linear
- (ii) MOLECULE TYPE: peptide
- (vi) ORIGINAL SOURCE:
  - (A) ORGANISM: Hirudunaria manillensis
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:4:

Val Ser Tyr Thr Asp Cys Thr Glu Ser Gly Gln Asn Tyr Cys Leu Cys
1 10 15

Val Gly Gly Asn Leu Cys Gly Gly Gly Lys His Cys Glu Met Asp Gly 20 25 30

Ser Gly Asn Lys Cys Val Asp Gly Glu Gly Thr Pro Lys Pro Lys Ser

Gln Thr Glu Gly Asp Phe Glu Glu Ile Pro Asp Glu Asp Ile Leu Asn 50 55 60

## (2) INFORMATION FOR SEQ ID NO:5:

- (i) SEQUENCE CHARACTERISTICS:
  - (A) LENGTH: 64 amino acids
  - (B) TYPE: amino acid
  - (C) STRANDEDNESS: single
  - (D) TOPOLOGY: linear
- (ii) MOLECULE TYPE: peptide
- (vi) ORIGINAL SOURCE:
  - (A) ORGANISM: Hirudinaris manillensis
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:5:
- Val Ser Tyr Thr Asp Cys Thr Glu Ser Gly Gln Asn Tyr Cys Leu Cys
  1 10 15
- Val Gly Ser Asn Val Cys Gly Glu Gly Lys Asn Cys Gln Leu Ser Ser 20 25 30
- Ser Gly Asn Gln Cys Val His Gly Glu Gly Thr Pro Lys Pro Lys Ser 35 40 45
- Gln Thr Glu Gly Asp Phe Glu Glu Ile Pro Asp Glu Asp Ile Leu Asn 50 55 60